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Database :
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Perfect score:
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Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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420.096 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	93.5	96	208	259	259	281	308	308	1170	1170	1206	Score
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	1435	257	170	49	49	ភូភ	60	60	674	674	234	Query Match Length DB
	20	22	22	22	22	22	22	22	21	21	21	DB
	AAY49070	AAU03696	AAB95297	AAM31512	AAM18938	AAM32811 .	AAM33333	AAM19849	AAB03758	AAB42482	AAB03759	ID
7	PolC gene product	Group B Streptococ	Human protein sequ	Peptide #5549 enco	Peptide #5372 enco	Peptide #6848 enco	Peptide #7370 enco	Peptide #6283 enco	Human H37 amino ac	Human ORFX ORF2246	Human H37 amino ac	Description

Human H37 proteins with a Cdc7 activity regulatory subunit, controlling cell replication and cell proliferation, useful

ior in treating

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82.5	82.5	82.5	83	83	83.5	84	84	84	84	84	84	84.5	84.5	84.5	84.5	84.5	84.5	84.5	84.5	84.5	84.5	85	85	85	85	86	86	87	87.5	88	68	89	93.5	
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AAG81779	AAW20098	AAW20667	AAM41108	AAM39322	AAG21891	AAB56018	AAY76079	AAY69163	AAY79154	AAB56062	AAY76123	AAY57452	AAY57453	AAM39431	AAY32187	AAY49248	AAY49151	AAG82087	AAG81684	AAG60518	AAR75416	AAW02289	AAY70209	AAB55946	AAY76007	AAY07114	AAW81168		AAG82479	AAR13227	AAY19934	AAY19935	AAB31934	
s. epidermidis ope		H. pylori cytoplas	Human polypeptide	Human polypeptide	Arabidopsis thalia	Skin cell protein,	Murine protein kin	Amino acid sequenc	Mouse protein kina	Skin cell protein,	Murine RIP protein	Human transcriptio	Human transcriptio	Human polypeptide			mino acid seq		S. epidermidis ope	Arabidopsis thalia	Rat regucalcin, a	Mouse neuron restr	Murine TANGO 130 p	skin cell protein,		WO9904265 Seq ID N	Transcriptional re	lautus endol	S. epidermidis ope	el endogluc	B. burgdorferi ant	B. burgdorferi ant	Amino acid sequenc	

ALIGNMENTS

RESULT AAB03759 H37; human; Cdc7 regulatory subunit; cytostatic; proliferative; cancer; anti-proliferative; replication regulator; stem cell. 04-OCT-2000 (first entry) WPI; 2000-365580/31. N-PSDB; AAA53484. Arai K, Masai H; (NISC-) JAPAN SCI & TECHNOLOGY CORP. (ARAI/) ARAI K. (MASA/) MASAI H. 30-OCT-1998; 01-NOV-1999; 11-MAY-2000. WO200026250-A1 Homo sapiens. Human H37 amino acid sequence #2. AAB03759; AAB03759 standard; Protein; 234 AA \vdash 98JP-0311408 99WO-JP06076

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31-MAR-1999;
02-APR-1999;
05-APR-1999;
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                                                                                                                                                                                                                                             Human; open reading frame; ORFX; detection; cytostatic; hepatotropic; vulnerary; antipsoriatic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; osteopathic; antiparkinsonian; nootropic; neuroprotective; anticonvulsant; oteropathic; antiparthritic; immunosuppressant; cardiant; immunostimulant; thrombolytic; coagulant; vasotropic; antidabetic; hypotensive; dermatological; immunosuppressive; antiinflammatory; antiviral; antibacterial; antifungal; antirheumatic; antithyroid; antianaemic; gene therapy; cancer; proliferative disorder; hypertension; neurodegenerative disorder; osteoarthritis; graft vs host disease; cardiovascular disease; osteoarthritis; hypothyroidism; SCID; AIDS; cholesterol ester storage; systemic lupus erythematosus; infection; severe combined immunodeficiency; malaria; autoimmune disorder; asthma; allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound; bone damage; cartilage damage; antiinflammatory disease; coagulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancers and diseases due to abnormal production of stem cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human ORFX ORF2246 polypeptide sequence SEQ ID NO:4492.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB42482 standard; Protein; 674 AA
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                                                                          31-MAR-2000; 2000WO-US08621.
                                                                                                                                                       WO200058473-A2.
                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                  thrombosis; contraceptive.
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99US-0127607.
99US-0127636.
99US-0127728.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AX AAC74446 to AAC77606 encode the proteins given in AAB40237 to AAB43397, CC which represent the human ORFX open reading frames 1 to 3161. The ORFX cequences have activities such as: cytostatic; hepatotropic; vulnerary; cc satisporiatic; antiparkinsonian; nootropic; neuroprotective; costeopathic; antiparkinsonian; nootropic; neuroprotective; antidabetic; hypotensive; dermatological; antinunosuppressive; antidabetic; hypotensive; dermatological; antifungal; antifungal; antifungal; antifungal; antifungal; antifungal; antiparkinsonia; conditions associated with an ORFX-associated districtive; costeopathicitis; corrective; and nucleic acids may be used to treat cancers, covectors. The proteins and nucleic acids may be used to treat cancers, covectors. The proteins and nucleic acids may be used to treat cancers, corrective disorders, osteopathitis, collecterial of soft disease, cardiovascular disease, diabetes mellitus, hypertension, hypothyroidism, cholesterol ester storage, systemic lupus crythematosus, severe combined immunodeficiency (SCID), AIDS, vital, collecterial or fungal infection, malaria, autoimmune disorders, asthma, collective, aplastic anaemia, burns, wounds, bone and cartilage damage, coccupiation; to inhibit thrombosis; and as a contraceptive.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel nucleic acids and peptides derived from open reading frame X, useful for treating e.g. cancers, proliferative disorders, neurodegenerative disorders and cardiovascular disease \dot{}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-MAR-2000; 2000US-0540763
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
04-OCT-2000 (first entry)
                                            AAB03758;
                                                                                           AAB03758 standard; Protein; 674 AA
                                                                                                                                                                                                                                                                                             181 qkkkelyllkksstsvrdggkrvgsgaqktrtgrlkkpfvkvedmsq 227
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DB; AAC76691.
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227; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 674;
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H37; human; Cdc7 regulatory subunit; cytostatic; proliferative; cancer; anti-proliferative; replication regulator; stem cell.

Human H37 amino acid sequence #1.

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ID AAM19849
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KW Cerv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; Fig 5; 55pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human H37 proteins with a Cdc7 activity regulatory subunit, controlling cell replication and cell proliferation, useful cancers and diseases due to abnormal production of stem cel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                        Probe; human; microarray;
                                                                                             12-OCT-2001 (first entry)
                                                                                                                                                    AAM19849 standard; Protein; 60
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(ARAI/) ARAI K.
(MASA/) MASAI H.
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                           cervical cancer.
                                                                                                                          AAM19849;
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                                                                   encoded by probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to abnormal production of stem cells
                                        gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                for measuring cervical gene expression
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                                         cervical
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                                        epithelial cell;
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                                                                                                                                                                                                                                                                                                                                                                      Peptide #7370 encoded by probe for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 27; SEQ ID No 24675; 487pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human genome-derived single exon nucleic acid probes useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Penn
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03-AUG-2000;
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26-MAY-2000;
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04-FEB-2000;
26-MAY-2000;
                                                                             30-JAN-2001;
                                                                                                                                  09-AUG-2001.
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                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                           genetic disorder
                                                                                                                                                                                                                                                                                                                    Probe; microarray; human; placenta; antenatal diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAM33333 standard; Protein;
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27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60 AA;
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2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0236359.
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2000US-0180312
2000US-0207456
                                                                             2001WO-US00663
                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in human cervical epithelial cells
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Pred. No. 8.8e-24;
0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                      measuring placental gene
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03-AUG-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
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26-MAY-2000; 2000US-0508408.

30-JUN-2000; 2000US-0508408.

03-AUG-2000; 2000US-0532366.

21-SEP-2000; 2000US-0234687.

21-SEP-2000; 2000US-0236359.

04-OCT-2000; 2000GB-0024263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 27; SEQ ID No 33602; 654pp; English.
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Human genome-derived single exon nucleic acid probes useful analyzing gene expression in human placenta \dot{\ }
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                                                                                   WPI; 2001-488897/53.
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2000US-0632366.
2000US-0234687.
2000US-0236359.
2000US-0024263.
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                                                                                                                                        Chen W,
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AAM18938
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26-MAY-2000; 2000US-0207456.

30-UUN-2000; 2000US-0608408.

03-AUG-2000; 2000US-0632366.

21-SEP-2000; 2000US-0234687.

27-SEP-2000; 2000US-0236359.

04-0CT-2000; 2000GB-0024263.
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The present invention relates to human single exon nucleic acid probes (SENP: see AAI10068-AAI28459). The present sequence is a peptide encoded by one such probe. The SENPs are derived from human HeLa cells. The SENPs can be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp. wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide #5372 encoded by probe for measuring cervical gene expression.
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                                                                                                                                                                                                                              Human genome-derived single exon nucleic acid probes useful i
analyzing gene expression in human cervical epithelial cells
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pred. No. 4.1e-21;
                                                                                                                                                                                                                                                                                                                    Rank DR;
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Query Match

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26-MAY-2000; 2000US-0207456.

30-JUN-2000; 2000US-0632366.

03-AUG-2000; 2000US-0632366.

21-SEP-2000; 2000US-0234687.

27-SEP-2000; 2000US-0236559.

04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1512
                                                                                                                         The present invention relates to single exon nucleic acid probes (SEMP: see AA.331315-AA.157546). The present sequence is a peptide encoded by one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of
                                                                                                                                                                                                                                                                  Penn
                                                                                                                                                                                          Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe; microarray; human; placenta; antenatal diagnosis;
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                     25 NRPSLKSLKTDNRPEKSKCKPLWGKVFYLDLPSVTISEKLQKDIKDLGG 73
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27;
                                                                                                                   genetic disorders.
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                                          49; Conserv
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                                                                                                                                                                                         SEQ ID No 31781; 654pp; English.
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ilarity 100.0%;
Conservative
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                                         Score 259; DB 22;
Pred. No. 5.8e-19;
0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 259;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .8e-19;
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                                                            Length 49;
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                                          Indels
 49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                          0;
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RESULT

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ω 5

kcqknspgark-

21 KNEKNRPSLKSLKTDNRPEKSKCKPLWGKVFYLDLPSVTISEKLQKDIKDLGGRVEEFLS 80

hpfsgksfyldlpagknlqfltgaiqqlggviegfls

Matches Query Match

Local

Similarity 36.4

17.2%; 36.4%;

Score 208; DB 22; Pred. No. 4.9e-13; 2; Mismatches 44;

Length 170;

30;

Gaps

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AAB95297
XX
AC AABS
AC COMMINICATION
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AC COMMINICATION
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AC COMMINICATION
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                                                             CC sequence and an oligonucleotide comprising a sequence complementary to a cc polynucleotide which comprises a 3'-end sequence, where the oligonucleotide which comprises at 1-sat 15 nucleotides and the combination of cc the 5'-end sequence3'-end sequence is selected from those defined in cc the specification. The primer sets can be used in antisense therapy and cc in gene therapy. The primer sets can be used in antisense therapy and cc particularly full-length cDNAs. The primers are also useful for the cc particularly full-length cDNAs. The primers are also useful for the cc the full-length cDNAs. The primers are also useful for the cc the full-length cDNAs. The primers allow obtaining of the full-length cDNAs easily without any specialised methods. AAH0316 to AAH13623 and CahH3633 to AAH18742 represent human cDNA sequences; AAB92446 to CC AAB95893 represent thuman amino acid sequences; AAB9246 to CC CAB95893 represent thuman amino acid sequences; AAB9246 to CC CAB95893 represent invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes primer sets for synthesising 5602 full-length cDNAs defined in the specification. Where a primer set comprises: (a) an oligo-off primer and an oligonucleotide complementary to the complementary strand of a polynucleotide which comprises one of the 5602 nucleotide sequences defined in the specification, where the oligonucleotide comprises at least 15 nucleotides; or (b) a combination of an oligonucleotide comprising a sequence complementary to the complementary strand of a polynucleotide which comprises a 5'-end
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer sets for synthesizing polynucleotides, particularly the 5602 full-length cDNAs defined in the specification, and for the detection and/or diagnosis of the abnormality of the proteins encoded by the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-AUG-1999; 99JP-0300253.
11-JAN-2000; 2000JP-0118776.
02-MAY-2000; 2000JP-0183767.
09-JUN-2000; 2000JP-0241899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ota T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JUL-2000; 2000EP-0116126
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human protein sequence SEQ ID NO:17525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAB95297;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB95297 standard; Protein;
       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   full-length cDNAs -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8; SEQ ID 17525;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isogai T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sugiyama T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
       ξ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detection; diagnosis; antisense therapy; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99JP-0248036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hikawa T, Hayashi K,
Wakamatsu A, Nagai I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2537pp +
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         170 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CD ROM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nagai K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saito K,
K, Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ξ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Yamamoto
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RESULT 10
AAMU369
XX
AC AAMU369
XX
AC AAMU369
XX
AC AAMU369
XX
AC Group 1
XX
Group 1
XW
Group 1
XW
Group 1
XW
WCapsid
XX
Strept
XX
WCapsid
XX
VCAPSID
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                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAU03601-AAU03722 represent Group B Streptococcus (Streptococcus agalactiae) amino acid sequences of the invention. S. agalactiae is an encapsulated bacterium which is a major pathogen of humans causing sepsis and meningitis in neonates as well as adults. The S. agalactiae antigenic polypeptides are used to vaccinate against Group B Streptococcus infections, particularly to prevent infection in new born children arising from the maternal genital tract. An immunogenic composition is useful in the preparation of a medicament for the treatment or prophylaxis of Group B Streptococcus infection. The invention does not have the disadvantages of varied response rate associated with prior art capsid polysaccharide vaccination against Group B Streptococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Fig 1; 178pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polypeptides derived from Streptococcus agalactiae are useful to provide detection of, and vaccination against, Group B Streptococcus infections, particularly to prevent infection in neonatals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-SEP-2000; 2000WO-GB03437.
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meningitis; neonate; antigenic; vaccine; infection; genital tract;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Group B Streptococcus antigenic protein, ID-173.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Le Page RWF, Wells JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MICR-) MICROBIAL TECHNICS LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200132882-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              capsid polysaccharide vaccination.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU03696 standard; Protein; 257 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              136 LSRGKLLVEKAIKDHDFIPSNSILSNALSWG 166
81 ylkqfykalqgyaksngvlelivepfddyqlftssgvpsnqgndnliedftssgyhhdgl 140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92 EAK-----FAQTLGRISPVPSPESAYTAETTSPHPSHDG------SSFKSPDT 133
                                                                                                                             23 flqtpemakllekrgydvrylgyqvenklelis--lsyimpvtggfqmkidsgpvhsnsk 80
                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001-316444/33.
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                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                          257 AA;
                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99GB-0021125.
                                                                                                                                                                                                                                                                                                     8.0%;
23.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hanniffy SB;
                                                                                                                                                                                                                                                                      34; Mismatches
                                                                                                                                                                                                                                                                                                     Score 96;
Pred. No.
                                                                                                                                                                                                                                                                                                                                    DB 22; Length 257;
                                                                                                                                                                                                                                                                                                     .18;
                                                                                                                                                                                                                                                                      54; Indels
                                                                                                                                                                                                                                                                  60; Gaps
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RESULT 11
AXY49070
ID AAY49070
XX AAY490
XX O5-JAN
XX Gram po
KW G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                δÃ
                                                              This is the PolC gene product Pol III-l of Staphylococcus aureus. The CC invention relates to a number of isolated DNA molecules from Gram CC positive bacterium, corresponding to dnaE (AAZ31001), and x (AAZ31002), and dnaB (AAZ31003). The PolC, dnaN and dnac genes (AAZ31002), and dnaB (AAZ31003). The PolC, dnaN and dnac genes (AAZ31002), and dnaB (AAZ31003). The PolC, dnaN and dnac genes (AAZ31002), and dnaB (AAZ31003). The polC, dnaN and dnac genes (AAZ31002), and dnaB (AAZ31003). The polC, dnaN and dnac genes (AAZ31002), and dnaB corresponds to the clapha subunit of the Escherichia coli, DNA polymerase III holdenzyme, complex corresponds to the gamma and tau subunits, and dnaB corresponds to the helicase. The alpha subunit is the actual DNA polymerase, the gamma complex forms the clamp loader and tau is a "glue protein". DnaX encodes complex forms the clamp loader and tau is a "glue protein". DnaX encodes complex forms and Tau, Tau is the product of the full gene, while gamma is the product of the first two thirds of the gene. The DNA sequences of the clavelous parts of the gene. The DNA sequences of the clavelous parts of the gene The DNA promote DNA creplication by acting on various parts of the gram positive bacterial DNA polymerase holoenzyme. The products and methods of the invention can be used for identifying pharmacological agents or lead compounds for agents or replication. The agents identified can be used as antibiotics.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gram positive bacteria; dnaE; dnaX; dnaB; PolC; dnaN; dnaG; helicase; alpha subunit; DNA polymerase III holoenzyme; gamma subunit; tau subunit; clamp loader; glue protein; replication; antibiotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated dnaE, dnaX and dnaB genes from Gram positive bacteria, used to develop screening assays for identifying antibiotic compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-590685/50.
N-PSDB; AAZ31004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JAN-1998;
22-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Key Location/Qualifiers Misc-difference 1033..1047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-JAN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY49070;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 25-30; 132pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-JUL-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09937661-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Staphylococcus aureus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PolC gene product Pol III-L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYRQ ) UNIV ROCKEFELLER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              141 ttgftgkylswhyvknlegvtsetllssfsktgralvkkamsfglkvrvl------k 191
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    192 rdelhlfkeitts 204
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     134 VCLSRGKLLVEKAIKDHDFIPSNSILSN------ALSWGVKILHIDDIRYYIEQK 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zhang D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0074522.
98US-0093727.
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Query Match

Best Local Similarity 25.1 Matches 57; Conservative

25.1%; -- 31;

Pred. No. 3.8; 1; Mismatches

;88

Indels

51;

Gaps

밁 δÕ В δÃ 8 δÃ В

967 nfsgeyqpn----ahnytkvlfgedkvfragtigtvaektafgyvk 1008

RYYIEQKKKELYLLKKSSTSVRDGGKRV-----GSGAQKTRTGRLK 216

915 seffndgsvgsgfdlpdktcetcgapl----ikegqdipfekflgfk---gdkvpdi-dl 966

-----DGSSFKSPDTVCLSRGKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDDI 175

858 ---gfaviylisqrlvkkslddgylvgsrgsvgssfvatmteitevnplpphyicpnckt 914

EFLSKDISYLISN---KKEAKFAQTLGRISPVPSPESAYTAETT--SPHPSH-----

123

77

805 vpikdelytprmeganeeirelsyanarklyge----dlpqivi-drlekelksiign-- 857

18 IQVKNEKNRPSLKSLKTDNRP-EKSKCKPLWGKVFYLDLPSVTISEKLQKDIKDLGGRVE 76

bacterial infection.

dnaE; Gram positive bacteria; polC; dnaE; holA; holB; dnaX; dnaN; ssb; dnaG; dnaB; antibiotic; replication; cell growth; cell death;

Amino acid sequence of a partial polC polypeptide.

15-MAY-2001 AAB31934;

(first entry)

AAB31934 standard;

Protein;

1435

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ARBSULT 1
ARAB31934
IID ARAB3
XX ARB31934
AC ARB3
XX ARB31934
AC ARB3
XX ARB3
AC ARB3
XX ARB3
AC ARB3

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     O'Donnell ME,
                                                                                                       29-JUL-1999;
                                                                                                                                                      28-JUL-2000; 2000WO-US20666.
                                                                                                                                                                                                         08-FEB-2001
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                                                   (UYRQ ) UNIV ROCKEFELLER
     Bruck I,
                                                                                                       99US-0146178
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1150
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1259
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1152
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                                                                                                                                                                                                                                                                                                                                                     /note- "encoded by
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Whipple
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AAY19935
ID AAY199
         PARRES PA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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                                                                                        03-SEP-1997;
20-JUN-1997;
22-JUL-1997;
22-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence represents a partial polC polypeptide. The specification describes DNA molecules from gram positive bacteria, which comprise a coding region from a polC, dnaE, holA, holB, dnaX, dnaN, ssb, dnaG or a dnaB gene. These sequences encode proteins that replicate the chromosome of Gram positive bacteria. They are used for sequencing and amplification of DNA and in drug discovery to identify compounds which have antibiotic activity through interference with replication. They are used in methods for identifying compounds that are active at the level of DNA replication and result in arrest of cell growth or cell death of bacteria to treat bacterial infections in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated DNA molecule from a Gram positive bacterium encoding DNA replication proteins used to identify compounds which have antibiotic activity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B. burgdorferi antigenic protein, t742.aa
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                                                                                                                                                                                                                                                                                                         30-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                        WO9859071-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                 Borrelia burgdorferi.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antigenic protein; vaccine; Lyme disease; infection; detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JUL-1999
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(HUMA-) HUMAN GENOME SCI INC
(MEDI-) MEDIMMUNE INC.
                                                                                                                                                                                                                                                   18-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           967 nfsgeyqpn----ahnytkvlfgedkvfragtigtvaektafgyvk 1008
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       176 RYYIEQKKKELYLLKKSSTSVRDGGKRV-----GSGAQKTRTGRLK 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18 IQVKNEKNRPSLKSLKTDNRP-EKSKCKPLWGKVFYLDLPSVTISEKLQKDIKDLGGRVE : :|:| : | : | : | : | : | : | : |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                seffndgsvgsgfdlpdktcetcgapl----ikegqdipfekflgfk---gdkvpdi-dl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                           97US-0057483.
97US-0050359.
97US-0053344.
97US-0053377.
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25.1%;
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Pred. No. 3
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RESULT 14
AAX19934
ID AAX199
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CS Borrel
XX Aorel
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Best Local
                         03-SEP-1997;
20-JUN-1997;
22-JUL-1997;
22-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This sequence represents a Borrelia burgdorferi (Bb) protein of the invention, which is suitable for use in a vaccine. The Bb polypeptides can be used in vaccines for eliciting protective antibodies to members of the Borrelia genus, particularly for the use against Lyme disease in humans and animals. They can be used for preventing or attenuating an infection caused by a member of the Borrelia genus. The products can also be used for detection of members of the Borrelia genus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New isolated Borrelia burgdorferi nucleic acids - used to develop products for the diagnosis, prevention and treatment of diseases caused by Borrelia, particularly Lyme disease
                                                                                                                                                                                                                                                                                                                                                  B. burgdorferi antigenic protein, f742.aa.
                                                                                                                                                                               30-DEC-1998.
                                                                                                                                                                                                                       WO9859071-A1
                                                                                                                                                                                                                                                               Borrelia burgdorferi
                                                                                                                                                                                                                                                                                                          Antigenic protein; vaccine; Lyme disease; infection; detection.
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                                                                                                                                    18-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAY19934 standard; Protein; 1119 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22 NEKNRPSLKSLKTDNRPEKSKCKP---LWGKVF--YLDLPSVTISEKLQKDIKDLG---- 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FVKVEDMSQSPAVH 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             eifqldkedkkpqyledlkskvhsikpidlentksr-qqaikdlneflknnpndaqaskt 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ----TAETTSPHPSHD-GSSFKSPDTVCLSRGKLLVEKAIKD-HDFI---PSNSILSNA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nnnnttslkkissnsqkeselsppsqtiigkiyrpy----sylikkelyeilddintgrv 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lagankighledlkskvhsikpidlentksrggaikdlneflknnpndagasktlagank 451
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LSWGVKILHIDDIRYYIEQ~KKKELYLLKKSSTSVRDGG---KRVGSGAQKTRTGRLKKP 218
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58; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Erwin AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1087 AA;
                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                         97US-0057483.
97US-0050359.
97US-0053344.
97US-0053377.
                                                                                                                                    98WO-US12718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         7.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hanson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 89;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches 101; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lathigra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 20; Length 1087;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Qy
                                                                   Cleavage-site
                                                                                                                                                                                                                                                               Cellulase activity; detergent.
                                                                                                                                                                                                                                                                                                          Novel endoglucanase
                                                                                                                                                                                                                                                                                                                                                       14-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                 AAR13227;
                           WO9110732-A
                                                                                                                                                                                                                    Bacillus spp. NCIMB 40250.
                                                                                                                                                        Peptide
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(first entry)

/label- mature endoglucanase 31..32

Location/Qualifiers /label signal sequence

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Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This sequence represents a Borrelia burgdorferi (Bb) protein of the invention, which is suitable for use in a vaccine. The Bb polypeptides can be used in vaccines for eliciting protective antibodies to members of the Borrelia genus, particularly for the use against Lyme disease in humans and animals. They can be used for preventing or attenuating an infection caused by a member of the Borrelia genus. The products can also be used for detection of members of the Borrelia genus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Choi GH, Erwin AL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 12; Page 124-125; 275pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New isolated Borrelia burgdorferi nucleic acids - used to develop products for the diagnosis, prevention and treatment of diseases caused by Borrelia, particularly Lyme disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-189980/16.
N-PSDB; AAX61631.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HUMA-) HUMAN GENOME SCI INC (MEDI-) MEDIMMUNE INC.
AAR13227 standard; Protein; 700 AA.
                                                                                                                                                      484 iqhledlksk--vh 495
                                                                                                                                                                                                                                                                               424 laqankiqhledlkskvhsikpidlentksrqqaikdlneflknnpndaqasktlaqank 483
                                                                                                                                                                                                                                                                                                                                                163
                                                                                                                                                                                                                                                                                                                                                                                                       365 eifqldkedkkpqyledlkskvhsikpidlentksr-qqaikdlneflknnpndaqaskt 423
                                                                                                                                                                                                                    219 FVKVEDMSQSPAVH 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    305 tigknrikelikkgisnkfqkvneliensknkeasnllitlikkdiepnlinipkdpykk 364
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             249 nnnnttslkkissnsqkeselsppsqtiigkiyrpy----sylikkelyeilddintgrv 304
                                                              15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           73 ----GRYEEFLSKDIS-----YLISNKKEAKFAQTL-----GRISP--VPSPESAY-- 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22 NEKNRPSLKSLKTDNRPEKSKCKP---LWGKVF--YLDLPSVTISEKLQKDIKDLG----
                                                                                                                                                                                                                                                                                                                                             LSWGVKILHIDDIRYYIEQ-KKKELYLLKKSSTSVRDGG---KRVGSGAQKTRTGRLKKP 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ----TAETTSPHPSHD-GSSFKSPDTVCLSRGKLLVEKAIKD-HDFI---PSNSILSNA 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity 58; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1119 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hanson MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 89; DB 20; Length 1119;
Pred. No. 7.6;
15; Mismatches 101; Indels 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lathigra R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 7.3%; Score 88; DB 12; Length 700; Best Local Similarity 23.9%; Pred. No. 4.8; Matches 54; Conservative 35; Mismatches 83; Indels 54; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The enzyme is produced by a strain of Bacillus spp. NCIMB 40250 and exhibits an endoglucanase activity of at least 10 (pref. at least 25) carboxymethyl cellulose (CMC) endoase units per mg total protein under alkaline conditions. It is especially useful as a cellulolytic agent and has been found to be more stable during washing (60 mins. at 40 deg.) in the presence of conventional detergents than a commercial cellulase preparation. It may also show increased storage stability in liq. detergents contg. proteases. The sequence was deduced from the DNA (AAQ13001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; Page 80; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Enzyme exhibiting cellulase activity from Bacillus sp. - is an endo-glucanase, esp. useful for harshness redn. of cotton-contg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; AAQ13001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jorgensen PL, Schulein M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (NOVO ) NOVO NORDISK A/S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-JAN-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUL-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    See also AAR13228 and AAR13229.
                                                                                                                                                                         239 --idlskavknvd--phaeifgpalygfgaylslqdapdwpslqgnyswfidyyldqmkn 294
                                                                                                                                                                                                                                                            185 mdeevnflvnrygnaststgikays--ldnepalwset---hprihpeqlqaaelvaks- 238
                                                                                             295 ahtqngkrlldvldvhwypeaqgggqrivfggagnidtqkarvqap 340
                                                                                                                      185 ELYLLKKSSTSVRD-------GGKRV----GSGAQKTRTGRLKKP 218
                                                                                                                                                                                                                                                                                                                                         142 vsrdkngpvdes-----etapsprwdkvefaknapf----slqpdlnd--gqv--y 184
                                                                                                                                                                                                                      139 GKLLVEKAIKDHDFIPSNSILSNALSWGVKILHIDD------IRYYIEQKKK 184
                                                                                                                                                                                                                                                                                                    79 LSKDISYLISNKKEAKFAQTLGRISPVPSPESAYTAETTSPHPSHDGSSFKSPDTVCLSR 138
                                                                                                                                                                                                                                                                                                                                                                                   20 VKNEKNRPSLKSLKTDNRPEKSKCKPLWGKV-FYLDLPSVTISEKLQKDIKDLGGRVEEF 78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             700 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hansen C;
                                                                                                                                                                                                                                                                                                                                                                                                                             12;
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